D. Clark

PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/08/999,690

DATE: 01/06/99

TIME: 16:38:00

RECEIVED APR 21 2000

TECH CENTER 1600/2900

INPUT SET: S30594.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1 2		SEQUENCE LISTING ENTERED					
3 4	(1) General Information:						
5 6 7 8	(i)	APPLICANT: Gunzburg, Walter H. Winder, David Saller, Robert M.					
9 10 11	(ii)	TITLE OF INVENTION: VECTORS CARRYING THERAPEUTIC GENES ENCODING ANTIMICROBIAL PEPTIDES FOR GENE THERAPY					
12 13	(iii)	NUMBER OF SEQUENCES: 12					
14 15 16 17 18 19 20	(iv)	CORRESPONDENCE ADDRESS: (A) ADDRESSEE: HAMILTON, BROOK, SMITH & REYNOLDS, P.C. (B) STREET: TWO MILITIA DRIVE (C) CITY: LEXINGTON (D) STATE: MASSACHUSETTS (E) COUNTRY: USA (F) ZIP: 02173					
22 23 24 25 26 27	(V)	COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.30					
28 29 30 31 32	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: US 08/999,690 (B) FILING DATE: 08-SEP-1997 (C) CLASSIFICATION:					
33 34 35 36	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: PCT/EP96/01001 (B) FILING DATE: 08-MAR-1996					
37 38 39 40	(vii)	PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: DK 0243/95 (B) FILING DATE: 09-MAR-1995					
41 42 43 44	(Viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: Brook, David E. (B) REGISTRATION NUMBER: 22,592 (C) REFERENCE/DOCKET NUMBER: GSF97-03A					
45 46	(ix)	TELECOMMUNICATION INFORMATION:					

RAW SEQUENCE LISTING PATENT APPLICATION US/08/999,690

TIME: 16:38:01

DATE: 01/06/99

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```
47
               (A) TELEPHONE: (781) 861-6240
48
               (B) TELEFAX: (781) 861-9540
49
50
    (2) INFORMATION FOR SEQ ID NO:1:
51
52
          (i) SEQUENCE CHARACTERISTICS:
53
54
               (A) LENGTH: 26 amino acids
55
               (B) TYPE: amino acid
56
               (C) STRANDEDNESS: unknown
               (D) TOPOLOGY: unknown
57
58
59
         (ii) MOLECULE TYPE: protein
60
61
62
63
64
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
65
         Gly Ile Gly Ala Val Leu Lys Val Leu Thr Thr Gly Leu Pro Ala Leu
66
67
                                               10
68
69
         Ile Ser Trp Ile Lys Arg Lys Arg Gln Gln
70
71
72
    (2) INFORMATION FOR SEQ ID NO:2:
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74
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76
               (B) TYPE: amino acid
               (C) STRANDEDNESS: unknown
77
78
               (D) TOPOLOGY: unknown
79
         (ii) MOLECULE TYPE: protein
80
81
82
83
84
85
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86
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87
88
89
90
         Ile Ser Trp Ile Gly Gly Gly Gly Gly
91
92
93
    (2) INFORMATION FOR SEQ ID NO:3:
94
95
          (i) SEQUENCE CHARACTERISTICS:
               (A) LENGTH: 25 base pairs
96
97
               (B) TYPE: nucleic acid
98
               (C) STRANDEDNESS: single
99
               (D) TOPOLOGY: linear
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PAGE: 3

RAW SEQUENCE LISTING PATENT APPLICATION US/08/999,690

DATE: 01/06/99 TIME: 16:38:02

INPUT SET: S30594.raw (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA" RECEIVED (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: TECH CENTER 1610 EXTEND TATGACGTCT CGTTAGAACG CGGCT (2) INFORMATION FOR SEQ ID NO:4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: GGCAGATCTT AAATGTATCA TGCAAT (2) INFORMATION FOR SEQ ID NO:5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: GACGTC (2) INFORMATION FOR SEQ ID NO:6: (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 6 base pairs

(C) STRANDEDNESS: single

(B) TYPE: nucleic acid

PAGE: 4

RAW SEQUENCE LISTING PATENT APPLICATION US/08/999,690

DATE: 01/06/99 TIME: 16:38:03

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153			(D) TOPOLOGY: linear	
154		, , , , ,	MOTEGUE MADE: other mucleic acid	
155		(11)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"	
156 157			(A) DESCRIPTION: /desc = DNA	
158				
159				
160			GROUPHOR REGORDSTON. GEO. ID NO. C.	
161		(X1)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
162		nam		
163	AGA	rcr		6
164		T.V.	DUNETON HOD GEO TO NO. 7	
165	(2)	INFO	RMATION FOR SEQ ID NO:7:	
166			anauruan autho amporantaa	
167		(1)	SEQUENCE CHARACTERISTICS:	
168			(A) LENGTH: 28 base pairs	
169			(B) TYPE: nucleic acid	
170	•		(C) STRANDEDNESS: single	
171			(D) TOPOLOGY: linear	
172				
173		(11)	MOLECULE TYPE: other nucleic acid	
174			(A) DESCRIPTION: /desc = "DNA"	
175				
176				
177				
178				
179		(XI)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
180				
181	ATA	BACGTO	CA AGGAAGGAAG CGATCGGA	28
182				
183	(2)	TNEO	RMATION FOR SEQ ID NO:8:	
184				
185		(1)	SEQUENCE CHARACTERISTICS:	
186			(A) LENGTH: 28 base pairs	
187			(B) TYPE: nucleic acid	
188			(C) STRANDEDNESS: single	
189			(D) TOPOLOGY: linear	
190				
191		(11)	MOLECULE TYPE: other nucleic acid	
192			(A) DESCRIPTION: /desc = "DNA"	
193				
194				
195				
196				
197		(Xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:	
198				
199	TAT	GGATC	CA ACCCTGTTGC CTCTTACG	28
200				
201	(2)	INFO	RMATION FOR SEQ ID NO:9:	
202				
203		(i)	SEQUENCE CHARACTERISTICS:	
204			(A) LENGTH: 35 base pairs	
205			(B) TYPE: nucleic acid	

RAW SEQUENCE LISTING PATENT APPLICATION US/08/999,690

DATE: 01/06/99 TIME: 16:38:04

INPUT SET: S30594.raw

206 207		(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
208			
209	(ii)	MOLECULE TYPE: other nucleic acid	
210		(A) DESCRIPTION: /desc = "DNA"	
211			
212	•		
213			
214			
215	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:	
216			
217	TCTTACAT	PCT ATGCGGGAAT TGGAGCAGTT CTGAA	35
218			
219	(2) INFO	ORMATION FOR SEQ ID NO:10:	
220			
221	(i)	SEQUENCE CHARACTERISTICS:	
222		(A) LENGTH: 35 base pairs	
223		(B) TYPE: nucleic acid	
224		(C) STRANDEDNESS: single	
225		(D) TOPOLOGY: linear	
226		•	
227	(ii)	MOLECULE TYPE: other nucleic acid	
228		(A) DESCRIPTION: /desc = "DNA"	
229			
230			
231			
232			
233	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:	
234			
235	AACTGCTC	CCA ATTCCCGCAT AGATGTAAGA AATGT	35
236		•	
237	(2) INFO	DRMATION FOR SEQ ID NO:11:	·
238			
239	(i)	SEQUENCE CHARACTERISTICS:	
240		· · · · · · · · · · · · · · · · · · ·	•
241		(B) TYPE: nucleic acid	
242		(C) STRANDEDNESS: single	
243		(D) TOPOLOGY: linear	
244			
245	(ii)	MOLECULE TYPE: other nucleic acid	
246		(A) DESCRIPTION: /desc = "DNA"	
247			
248			
249		,	
250			
251	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:	•
252			
253	GACGTC		6
254			
255	(2) INFO	DRMATION FOR SEQ ID NO:12:	
256			
257	(i)	SEQUENCE CHARACTERISTICS:	
258		(A) LENGTH: 6 base pairs	

PAGE: 1

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/999,690

DATE: 01/06/99 TIME: 16:38:05

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Original Text